SEQUENCE LISTING

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	cct Pro															288	·	
	atc Ile															336		
	tat Tyr															384		
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Gly	Glu 690	Gln	Cys	Glu	Gly	Glu 695	Glu	Asp	Thr	Glu	Phe 700	Met	Thr	Pro	Ser
Ser 705	Arg	Pro	Leu	Arg	Pro 710	Leu	Asp	Thr	Ser	Gln 715	Ser	Ser	Arg	Ala	Cys 720
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Ile	Gln	Ser	Gln 740	Ala	Pro	Ser	Ile	Thr 745		Ser		Thr	Phe 750	Gly	Glu

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cat His 210												672
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Ser Gly Gly Ser Gly Ser Gly Leu Ile Gly Leu Met Lys Asp Ala Page 86

10

Phe Gln Pro His His His His His His Leu Ser Pro His Pro Pro

20

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Lys Val Val Arg Leu Cys Gln Asn Pro Lys Leu Ala Leu Lys Asn Ser 65 70 75 80

Pro Pro Tyr Ile Leu Asp Leu Leu Pro Asp Thr Tyr Gln His Leu Arg 85 90 95

Thr Ile Leu Ser Arg Tyr Glu Gly Lys Met Glu Thr Leu Gly Glu Asn 100 105 110

Glu Tyr Phe Arg Val Phe Met Glu Asn Leu Met Lys Lys Thr Lys Gln 115 120 125

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Asp Thr Phe Arg Ile Thr Lys Ala Asp Ala Ala Glu Phe Trp Arg Lys 180 185 190

Ala Phe Gly Glu Lys Thr Ile Val Pro Trp Lys Ser Phe Arg Gln Ala 195 200 205

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Phe Asp Ile Phe Thr Arg Leu Phe Gln Pro Trp Ser Ser Leu Leu Arg
245 250 255

Asn Trp Asn Ser Leu Ala Val Thr His Pro Gly Tyr Met Ala Phe Leu Page 87 260 265 270

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500 505 510

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Asn Glu Asp Asp Gly Tyr Asp Val Pro Lys Pro Pro Val Pro Ala Val 770 775 780

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gat t Asp (2208

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Phe	Gln	Pro 35	His	His	His	His	His 40	His	His	Leu	Ser	Pro 45	His	Pro	Pro
Gly	Thr 50	Val	Asp	Lys	Lys	Met 55	Val	Glu	Lys	Cys	Trp 60	Lys	Leu	Met	Asp
Lys 65	Val	Val	Arg	Leu	Cys 70	Gln	Asn	Pro	Lys	Leu 75	Ala	Leu	Lys	Asn	Ser 80
Pro	Pro	Tyr	Ile	Leu 85	Asp	Leu	Leu	Pro	Asp 90	Thr	Tyr	Gln	His	Leu 95	Arg
Thr	Ile	Leu	Ser 100	Arg	Tyr	Glu	Gly	Lys 105	Met	Glu	Thr	Leu	Gly 110	Glu	Asn
Glu	Tyr	Phe 115	Arg	Val	Phe	Met	Glu 120	Asn	Leu	Met	Lys	Lys 125	Thr	Lys	Gln
Thr	Ile 130	Ser	Leu	Phe	Lys	Glu 135	Gly	Lys	Glu	Arg	Met 140	Tyr	Glu	Glu	Asn
Ser 145	Gln	Pro	Arg	Arg	Asn 150	Leu	Thr	Lys	Leu	Ser 155	Leu	Ile	Phe	Ser	His 160
Met	Leu	Ala	Glu	Leu 165	Lys	Gly	Ile	Phe	Pro 170	Ser	Gly	Leu	Phe	Gln 175	Gly
Asp	Thr	Phe	Arg 180	Ile	Thr	Lys	Ala	Asp 185	Ala	Ala	Glu	Phe	Trp 190	Arg	Lys
Ala	Phe	Gly 195	Glu	Lys	Thr	Ile	Val 200	Pro	Trp	Lys	Ser	Phe 205	Arg	Gln	Ala
Leu	His 210	Glu	Val	His	Pro	Ile 215	Ser	Ser	Gly	Leu	Glu 220	Ala	Met	Ala	Leu
Lys 225	Ser	Thr	Ile	Asp	Leu 230	Thr	Cys	Asn	Asp	Tyr 235	Ile	Ser	Val	Phe	Glu 240
Phe	Asp	Ile	Phe	Thr 245	Arg	Leu	Phe	Gln	Pro 250	Trp	Ser	Ser	Leu	Leu 255	Arg

Asn	Trp	Asn	Ser 260	Leu	Ala	Val	Thr	His 265	Pro	Gly	Tyr	Met	Ala 270	Phe	Leu
Thr	Tyr	Asp 275	Glu	Val	Lys	Ala	Arg 280	Leu	Gln	Lys	Phe	Ile 285	His	Lys	Pro
Gly	Ser 290	Tyr	Ile	Phe	Arg	Leu 295	Ser	Cys	Thr	Arg	Leu 300	Gly	Gln	Trp	Ala
Ile 305	Gly	Tyr	Val	Thr	Ala 310	Asp	Gly	Asn	Ile	Leu 315	Gln	Thr	Ile	Pro	His 320
Asn	Lys	Pro	Leu	Phe 325	Gln	Ala	Leu	Ile	Asp 330	Gly	Phe	Arg	Glu	Gly 335	Phe
Tyr	Leu	Phe	Pro 340	Asp	Gly	Arg	Asn	Gln 345	Asn	Pro	Asp	Leu	Thr 350	Gly	Leu
Cys	Glu	Pro 355	Thr	Pro	Gln	Asp	His 360	Ile	Lys	Val	Thr	Gln 365	Glu	Gln	Tyr
Glu	Leu 370	Tyr	Cys	Glu	Met	Gly 375	Ser	Thr	Phe	Gln	Leu 380	Cys	Lys	Ile	Cys
Ala 385	Glu	Asn	Asp	Lys	Asp 390	Val	Lys	Ile	Glu	Pro 395	Cys	Gly	His	Leu	Met 400
Cys	Thr	Ser	Cys	Leu 405	Thr	Ser	Trp	Gln	Glu 410	Ser	Glu	Gly	Gln	Gly 415	Cys
Pro	Phe	Суѕ	Arg 420	Cys	Glu	Ile	Lys	Gly 425	Thr	Glu	Pro	Ile	Val 430	Val	Asp
Pro	Phe	Asp 435	Pro	Arg	Gly	Ser	Gly 440	Ser	Leu	Leu	Arg	Gln 445	Gly	Ala	Glu
Gly	Ala 450	Pro	Ser	Pro	Asn	Tyr 455	Asp	Asp	Asp	Asp	Asp 460	Glu	Arg	Ala	Asp
Asp 465	Thr	Leu	Phe	Met	Met 470	Lys	Glu	Leu	Ala	Gly 475	Ala	Lys	Val	Glu	Arg 480
Pro	Pro	Ser	Pro	Phe 485	Ser	Met	Ala	Pro	Gln 490	Ala	Ser	Leu	Pro	Pro 495	Val

Pro	Pro	Arg	Leu 500	Asp	Leu	Leu	Pro	Gln 505	Arg	Val	Cys	Val	Pro 510	Ser	Ser
Ala	Ser	Ala 515	Leu	Gly	Thr	Ala	Ser 520	Lys	Ala	Ala	Ser	Gly 525	Ser	Leu	His
Lys	Asp 530	Lys	Pro	Leu	Pro	Val 535	Pro	Pro	Thr	Leu	Arg 540	Asp	Leu	Pro	Pro
Pro 545	Pro	Pro	Pro	Asp	Arg 550	Pro	Tyr	Ser	Val	Gly 555	Ala	Glu	Ser	Arg	Pro 560
Gln	Arg	Arg	Pro	Leu 565	Pro	Cys	Thr	Pro	Gly 570	Asp	Cys	Pro	Ser	Arg 575	Asp
Lys	Leu	Pro	Pro 580	Val	Pro	Ser	Ser	Arg 585	Leu	Gly	Asp	Ser	Trp 590	Leu	Pro
Arg	Pro	Ile 595	Pro	Lys	Val	Pro	Val 600	Ser	Ala	Pro	Ser	Ser 605	Ser	Asp	Pro
Trp	Thr 610	Gly	Arg	Glu	Leu	Thr 615	Asn	Arg	His	Ser	Leu 620	Pro	Phe	Ser	Leu
Pro 625	Ser	Gln	Met	Glu	Pro 630	Arg	Pro	Asp	Val	Pro 635	Arg	Leu	Gly	Ser	Thr 640
Phe	Ser	Leu	Asp	Thr 645	Ser	Met	Ser	Met	Asn 650	Ser	Ser	Pro	Leu	Val 655	Gly
Pro	Glu	Cys	Asp 660	His	Pro	Lys	Ile	Lys 665	Pro	Ser	Ser	Ser	Ala 670	Asn	Ala
Ile	Tyr	Ser 675	Leu	Ala	Ala	Arg	Pro 680	Leu	Pro	Val	Pro	Lys 685	Leu	Pro	Pro
Gly	Glu 690	Gln	Cys	Glu	Gly	Glu 695	Glu	Asp	Thr	Glu	Tyr 700	Met	Thr	Pro	Ser
Ser 705	Arg	Pro	Leu	Arg	Pro 710	Leu	Asp	Thr	Ser	Gln 715	Ser	Ser	Arg	Ala	Cys 720
Asp	Cys	Asp	Gln	Gln 725	Ile	Asp	Ser	Cys	Thr 730	Tyr	Glu	Ala	Met	Tyr 735	Asn

Ile Gln Ser Gln Ala Pro Ser Ile Thr Glu Ser Ser Thr Phe Gly Glu 745 Gly Asn Leu Ala Ala Ala His Ala Asn Thr Gly Pro Glu Glu Ser Glu 760 Asn Glu Asp Asp Gly Phe Asp Val Pro Lys Pro Pro Val Pro Ala Val 775 Leu Ala Arg Arg Thr Leu Ser Asp Ile Ser Asn Ala Ser Ser Ser Phe 795 800 785 Gly Trp Leu Ser Leu Asp Gly Asp Pro Thr Thr Asn Val Thr Glu Gly 805 810 Ser Gln Val Pro Glu Arg Pro Pro Lys Pro Phe Pro Arg Arg Ile Asn 820 825 830 Ser Glu Arg Lys Ala Gly Ser Cys Gln Gln Gly Ser Gly Pro Ala Ala 835 840 845 Ser Ala Ala Thr Ala Ser Pro Gln Leu Ser Ser Glu Ile Glu Asn Leu 850 855 860 Met Ser Gln Gly Tyr Ser Tyr Gln Asp Ile Gln Lys Ala Leu Val Ile 865 Ala Gln Asn Asn Ile Glu Met Ala Lys Asn Ile Leu Arg Glu Phe Val 885 895 Ser Ile Ser Ser Pro Ala His Val Ala Thr 900 <210> 257 <211> 2721 <212> DNA <213> artificial <220> <223> c-Cbl Y700F/Y731F/Y774F triple mutant <220> <221> CDS <222> (1)..(2718) <400> 257

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		cac ctc agc ccc cac His Leu Ser Pro His 45	
		aag tgc tgg aag ctc Lys Cys Trp Lys Leu 60	
		aag ctg gcg cta aag Lys Leu Ala Leu Lys 75	
		gat acc tac cag cat Asp Thr Tyr Gln His 90	_
	Arg Tyr Glu Gly Lys	atg gag aca ctt gga Met Glu Thr Leu Gly 110	
		ttg atg aag aaa act Leu Met Lys Lys Thr 125	
		gaa aga atg tat gag Glu Arg Met Tyr Glu 140	
		ctg tcc ctc atc ttc Leu Ser Leu Ile Phe 155	
		cca agt gga ctc ttt Pro Ser Gly Leu Phe 170	
	Ile Thr Lys Ala Asp	gct gcg gaa ttt tgg Ala Ala Glu Phe Trp 190	
		tgg aag agc ttt cga Trp Lys Ser Phe Arg 205	
		ggg ctg gag gcc atg Gly Leu Glu Ala Met 220	
		gat tat att tcg gtt Asp Tyr Ile Ser Val 235	
		g ccc tgg tcc tct ttg n Pro Trp Ser Ser Leu Page 99	

	245	250	255
		cat cct ggc tac His Pro Gly Tyr 265	
		ctc cag aaa ttc Leu Gln Lys Phe	
-		tgt act cgt ctg Cys Thr Arg Leu 300	
		aac att ctc cag Asn Ile Leu Gln 315	
		att gat ggc ttc Ile Asp Gly Phe 330	
	Asp Gly Arg Asn	cag aat cct gat Gln Asn Pro Asp 345	
		atc aaa gtg acc Ile Lys Val Thr	
=		aca ttc caa cta Thr Phe Gln Leu 380	
		att gag ccc tgt Ile Glu Pro Cys 395	
		cag gaa tca gaa Gln Glu Ser Glu 410	
		ggt act gaa ccc Gly Thr Glu Pro 425	
		agc ctg ttg agg Ser Leu Leu Arg	
		gat gat gat gat Asp Asp Asp Asp 460	
		ttg gct ggt gcc Leu Ala Gly Ala 475	
		cca caa gct tcc Pro Gln Ala Ser Page 100	ctt ccc ccg gtg 1488 Leu Pro Pro Val

48	5	490	495
	p Leu Leu Pro G	ag cga gta tgt gtt ln Arg Val Cys Val 05	
		ag gct gct tct ggc ys Ala Ala Ser Gly 525	
_	-	cc aca ctt cga gat ro Thr Leu Arg Asp 540	
		ct gtt gga gca gaa er Val Gly Ala Glu 555	
	eu Pro Cys Thr Pr	ca ggc gac tgt ccc ro Gly Asp Cys Pro 570	
	l Pro Ser Ser A	gc ctt gga gac tca rg Leu Gly Asp Ser 85	
	-	ct gcc cca agt tcc er Ala Pro Ser Ser 605	
		gg cac tca ctt cca rg His Ser Leu Pro 620	
		at gtg cct agg ctc sp Val Pro Arg Leu 635	
· · · · · ·	r Ser Met Ser Me	tg aat agc agc cca et Asn Ser Ser Pro 650	• ••
	s Pro Lys Ile Ly	aa cct tcc tca tct ys Pro Ser Ser Ser 65	
		tt cct gtg cca aaa eu Pro Val Pro Lys 685	
		ac aca gag ttc atg sp Thr Glu Phe Met 700	
		ca tcc cag agt tca hr Ser Gln Ser Ser 715	
		gt acg ttt gaa gca ys Thr Phe Glu Ala Page 101	

725		730	735
		gag agc agc acc ttt Glu Ser Ser Thr Phe 750	
		act ggt ccc gag gag Thr Gly Pro Glu Glu 765	
		aag cca cct gtg ccg Lys Pro Pro Val Pro 780	
		tct aat gcc agc tcc Ser Asn Ala Ser Ser 795	
	Asp Gly Asp Pro	aca aca aat gtc act Thr Thr Asn Val Thr 810	
		cca ttc ccg cgg aga Pro Phe Pro Arg Arg 830	
		caa ggt agt ggt cct Gln Gly Ser Gly Pro 845	
		tcc agt gag atc gag Ser Ser Glu Ile Glu 860	
		atc cag aaa gct ttg Ile Gln Lys Ala Leu 875	_
	Glu Met Ala Lys	aac atc ctc cgg gaa Asn Ile Leu Arg Glu 890	_
tcc att tct tct cct Ser Ile Ser Ser Pro 900			2721
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<220> <223> Synthetic Co	nstruct		
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Phe	Gln	Pro 35	His	His	His	His	His 40	His	His	Leu	Ser	Pro 45	His	Pro	Pro
Gly	Thr 50	Val	Asp	Lys	Lys	Met 55	Val	Glu	Lys	Cys	Trp 60	Lys	Leu	Met	Asp
Lys 65	Val	Val	Arg	Leu	Cys 70	Gln	Asn	Pro	Lys	Leu 75	Ala	Leu	Lys	Asn	Ser 80
Pro	Pro	Tyr	Ile	Leu 85	Asp	Leu	Leu	Pro	Asp 90	Thr	Tyr	Gln	His	Leu 95	Arg
Thr	Ile	Leu	Ser 100	Arg	Tyr	Glu	Gly	Lys 105	Met	Glu	Thr	Leu	Gly 110	Glu	Asn
Glu	Tyr	Phe 115	Arg	Val	Phe	Met	Glu 120	Asn	Leu	Met	Lys	Lys 125	Thr	Lys	Gln
Thr	Ile 130	Ser	Leu	Phe	Lys	Glu 135	Gly	Lys	Glu	Arg	Met 140	Tyr	Glu	Glu	Asn
Ser 145	Gln	Pro	Arg	Arg	Asn 150	Leu	Thr	Lys	Leu	Ser 155	Leu	Ile	Phe	Ser	His 160
Met	Leu	Ala	Glu	Leu 165	Lys	Gly	Ile	Phe	Pro 170	Ser	Gly	Leu	Phe	Gln 175	Gly
Asp	Thr	Phe	Arg 180	Ile	Thr	Lys	Ala	Asp 185	Ala	Ala	Glu	Phe	Trp 190	Arg	Lys
Ala	Phe	Gly 195	Glu	Lys	Thr	Ile	Val 200	Pro	Trp	Lys	Ser	Phe 205	Arg	Gln	Ala
Leu	His 210	Glu	Val	His	Pro	Ile 215	Ser	Ser	Gly	Leu	Glu 220	Ala	Met	Ala	Leu
Lys 225	Ser	Thr	Ile	Asp	Leu 230	Thr	Cys	Asn	Asp	Tyr 235	Ile	Ser	Val	Phe	Glu 240
Phe	Asp	Ile	Phe	Thr 245	Arg	Leu	Phe	Gln	Pro 250	Trp	Ser	Ser	Leu	Leu 255	Arg

Asn Trp	Asn Ser 260	Leu Ala	Val	Thr	His 265	Pro	Gly	Tyr	Met	Ala 270	Phe	Leu
Thr Tyr	Asp Glu 275	Val Lys	Ala	Arg 280	Leu	Gln	Lys	Phe	Ile 285	His	Lys	Pro
Gly Ser 1	Tyr Ile	Phe Arg	Leu 295	Ser	Cys	Thr	Arg	Leu 300	Gly	Gln	Trp	Ala
Ile Gly '	Tyr Val	Thr Ala 310	Asp	Gly	Asn	Ile	Leu 315	Gln	Thr	Ile	Pro	His 320
Asn Lys	Pro Leu	Phe Gln 325	Ala	Leu	Ile	Asp 330	Gly	Phe	Arg	Glu	Gly 335	Phe
Tyr Leu	Phe Pro 340	Asp Gly	Arg	Asn	Gln 345	Asn	Pro	Asp	Leu	Thr 350	Gly	Leu
Cys Glu	Pro Thr 355	Pro Gln	Asp	His 360	Ile	Lys	Val	Thr	Gln 365	Glu	Gln	Tyr
Glu Leu '	Tyr Cys	Glu Met	Gly 375	Ser	Thr	Phe	Gln	Leu 380	Cys	Lys	Ile	Cys
Ala Glu . 385	Asn Asp	Lys Asp 390		Lys	Ile	Glu	Pro 395	Cys	Gly	His	Leu	Met 400
Cys Thr	Ser Cys	Leu Thr 405	Ser	Trp	Gln	Glu 410	Ser	Glu	Gly	Gln	Gly 415	Cys
Pro Phe	Cys Arg 420	Cys Glu	Ile	Lys	Gly 425	Thr	Glu	Pro	Ile	Val 430	Val	Asp
Pro Phe	Asp Pro 435	Arg Gly	Ser	Gly 440	Ser	Leu	Leu	Arg	Gln 445	Gly	Ala	Glu
Gly Ala 450	Pro Ser	Pro Asn	Tyr 455	Asp	Asp	Asp	Asp	Asp 460	Glu	Arg	Ala	Asp
Asp Thr 465	Leu Phe	Met Met 470	-	Glu	Leu	Ala	Gly 475	Ala	Lys	Val	Glu	Arg 480
Pro Pro	Ser Pro	Phe Ser 485	Met	Ala	Pro	Gln 490	Ala	Ser	Leu	Pro	Pro 495	Val

Pro P	ro	Arg	Leu 500	Asp	Leu	Leu	Pro	Gln 505	Arg	Val	Cys	Val	Pro 510	Ser	Ser
Ala S		Ala 515	Leu	Gly	Thr	Ala	Ser 520	Lys	Ala	Ala	Ser	Gly 525	Ser	Leu	His
Lys A 5	sp 30	Lys	Pro	Leu	Pro	Val 535	Pro	Pro	Thr	Leu	Arg 540	Asp	Leu	Pro	Pro
Pro P 545	ro	Pro	Pro	Asp	Arg 550	Pro	Tyr	Ser	Val	Gly 555	Ala	Glu	Ser	Arg	Pro 560
Gln A	rg	Arg	Pro	Leu 565	Pro	Cys	Thr	Pro	Gly 570	Asp	Cys	Pro	Ser	Arg 575	Asp
Lys L	eu	Pro	Pro 580	Val	Pro	Ser	Ser	Arg 585	Leu	Gly	Asp	Ser	Trp 590	Leu	Pro
Arg P	ro	Ile 595	Pro	Lys	Val	Pro	Val 600	Ser	Ala	Pro	Ser	Ser 605	Ser	Asp	Pro
Trp T	hr 510	Gly	Arg	Glu	Leu	Thr 615	Asn	Arg	His	Ser	Leu 620	Pro	Phe	Ser	Leu
Pro S 625	Ser	Gln	Met	Glu	Pro 630	Arg	Pro	Asp	Val	Pro 635	Arg	Leu	Gly	Ser	Thr 640
Phe S	Ser	Leu	Asp	Thr 645	Ser	Met	Ser	Met	Asn 650	Ser	Ser	Pro	Leu	Val 655	Gly
Pro G	lu	Cys	Asp 660	His	Pro	Lys	Ile	Lys 665	Pro	Ser	Ser	Ser	Ala 670	Asn	Ala
Ile T	ſyr	Ser 675	Leu	Ala	Ala	Arg	Pro 680	Leu	Pro	Val	Pro	Lys 685	Leu	Pro	Pro
Gly G	31u 590	Gln	Cys	Glu	Gly	Glu 695	Glu	Asp	Thr	Glu	Phe 700	Met	Thr	Pro	Ser
Ser <i>A</i> 705	Arg	Pro	Leu	Arg	Pro 710	Leu	Asp	Thr	Ser	Gln 715	Ser	Ser	Arg	Ala	Cys 720
Asp (Cys	Asp	Gln	Gln 725	Ile	Asp	Ser	Cys	Thr 730	Phe	Glu	Ala	Met	Туг 735	Asn

Ile Gl
n Ser Gl
n Ala Pro Ser Ile Thr Glu Ser Ser Thr Phe Gly Glu
 740 745 750

Gly Asn Leu Ala Ala Ala His Ala Asn Thr Gly Pro Glu Glu Ser Glu 755 760 765

Asn Glu Asp Asp Gly Phe Asp Val Pro Lys Pro Pro Val Pro Ala Val 770 780

Leu Ala Arg Arg Thr Leu Ser Asp Ile Ser Asn Ala Ser Ser Ser Phe 785 790 795 800

Gly Trp Leu Ser Leu Asp Gly Asp Pro Thr Thr Asn Val Thr Glu Gly 805 810 815

Ser Gln Val Pro Glu Arg Pro Pro Lys Pro Phe Pro Arg Arg Ile Asn 820 825 830

Ser Glu Arg Lys Ala Gly Ser Cys Gln Gln Gly Ser Gly Pro Ala Ala 835 840 845

Ser Ala Ala Thr Ala Ser Pro Gln Leu Ser Ser Glu Ile Glu Asn Leu 850 855 860

Met Ser Gln Gly Tyr Ser Tyr Gln Asp Ile Gln Lys Ala Leu Val Ile 865 870 875 880

Ala Gln Asn Asn Ile Glu Met Ala Lys Asn Ile Leu Arg Glu Phe Val 885 890 895

Ser Ile Ser Ser Pro Ala His Val Ala Thr 900 905

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<400> 259

atg gcc ggc aac gtg aag aag agc tct ggg gcc ggg ggc ggc acg ggc Page 106

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				ggt Gly												96	
				cac His												144	
				aag Lys												192	
				ttg Leu												240	
				tta Leu 85												288	
				aga Arg												336	
				gtg Val												384	
				ttc Phe												432	
				cga Arg												480	
		Ăla	Glu	cta Leu 165	Lys	Gly	Ile	Phe	Pro	Ser	Gly	Leu	Phe	Gln	Gly	528	
				att Ile											aaa Lys	576	
gct Ala	ttt Phe	ggg Gly 195	Glu	aag Lys	aca Thr	ata Ile	gtc Val 200	cct Pro	tgg Trp	aag Lys	agc Ser	ttt Phe 205	cga Arg	cag Gln	gct Ala	624	
cta Leu	cat His 210	Glu	gtg Val	cat His	ccc Pro	atc Ile 215	agt Ser	tct Ser	Gly	ctg Leu	gag Glu 220	gcc Ala	atg Met	gct Ala	ctg Leu	672	
						Thr					Ile				gaa Glu 240	720	
ttt	gac	atc	ttt	acc	cga	ctc	ttt	cag		tgg age		tct	ttg	ctc	agg	768	

Phe	Asp	Ile	Phe	Thr 245	Arg	Leu	Phe	Gln	Pro 250	Trp	Ser	Ser	Leu	Leu 255	Arg	
								cat His 265								816
								ctc Leu								864
								tgt Cys								912
								aac Asn								960
								att Ile								1008
								cag Gln 345								1056
								atc Ile								1104
								aca Thr								1152
								att Ile								1200
		Ser	Cys	Leu	Thr	Ser	Trp	cag Gln	Glu	Ser	Glu	Gly	Gln	Gly	Cys	1248
cct Pro	ttc Phe	tgc Cys	cga Arg 420	tgt Cys	gaa Glu	att Ile	aaa Lys	ggt Gly 425	act Thr	gaa Glu	ccc Pro	atc Ile	gtg Val 430	gta Val	gat Asp	1296
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Ala	Phe	Gly 195	Glu	Lys	Thr	Ile	Val 200	Pro	Trp	Lys	Ser	Phe 205	Arg	Gln	Ala
Leu	His 210	Glu	Val	His	Pro	Ile 215	Ser	Ser	Gly	Leu	Glu 220	Ala	Met	Ala	Leu
Lys 225	Ser	Thr	Ile	Asp	Leu 230	Thr	Cys	Asn	Asp	Туг 235	Ile	Ser	Val	Phe	Glu 240
Phe	Asp	Ile	Phe	Thr 245	Arg	Leu	Phe	Gln	Pro 250	Trp	Ser	Ser	Leu	Leu 255	Arg
Asn	Trp	Asn	Ser 260	Leu	Ala	Val	Thr	His 265	Pro	Gly	Tyr	Met	Ala 270	Phe	Leu
Thr	Tyr	Asp 275	Glu	Val	Lys	Ala	Arg 280	Leu	Gln	Lys	Phe	Ile 285	His	Lys	Pro
Gly	Ser 290	Tyr	Ile	Phe	Arg	Leu 295	Ser	Cys	Thr	Arg	Leu 300	Gly	Gln	Trp	Ala
Ile 305	Gly	Tyr	Val	Thr	Ala 310	Asp	Gly	Asn	Ile	Leu 315	Gln	Thr	Ile	Pro	His 320
Asn	Lys	Pro	Leu	Phe 325	Gln	Ala	Leu	Ile	Asp 330	Gly	Phe	Arg	Glu	Gly 335	Phe
Tyr	Leu	Phe	Pro 340	Asp	Gly	Arg	Asn	Gln 345	Asn	Pro	Asp	Leu	Thr 350	Gly	Leu
Cys	Glu	Pro 355	Thr	Pro	Gln	Asp	His 360	Ile	Lys	Val	Thr	Gln 365	Glu	Gln	Tyr
Glu	Leu 370	Tyr	Cys	Glu	Met	Gly 375	Ser	Thr	Phe	Gln	Leu 380	Cys	Lys	Ile	Cys
Ala 385	Glu	Asn	Asp	Lys	Asp 390	Val	Lys	Ile	Glu	Pro 395	Cys	Gly	His	Leu	Met 400
Cys	Thr	Ser	Суз	Leu 405	Thr	Ser	Trp	Gln	Glu 410	Ser	Glu	Gly	Gln	Gly 415	Cys
Pro	Phe	Cys	Arg 420	Суз	Glu	Ile	Lys	Gly 425	Thr	Glu	Pro	Ile	Val 430	Val	Asp

Pro Phe Asp Pro Arg Gly Ser Gly Ser Leu Leu Arg Gln Gly Ala Glu 435 440 445

Gly Ala Pro Ser Pro Asn Tyr Asp Asp Asp Asp Glu Arg Ala Asp 450 455 460

Asp Thr Leu Phe Met Met Lys Glu Leu Ala Gly Ala Lys Val Glu Arg 465 470 475 480

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<212> DNA

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